

Supplementary Figure 2

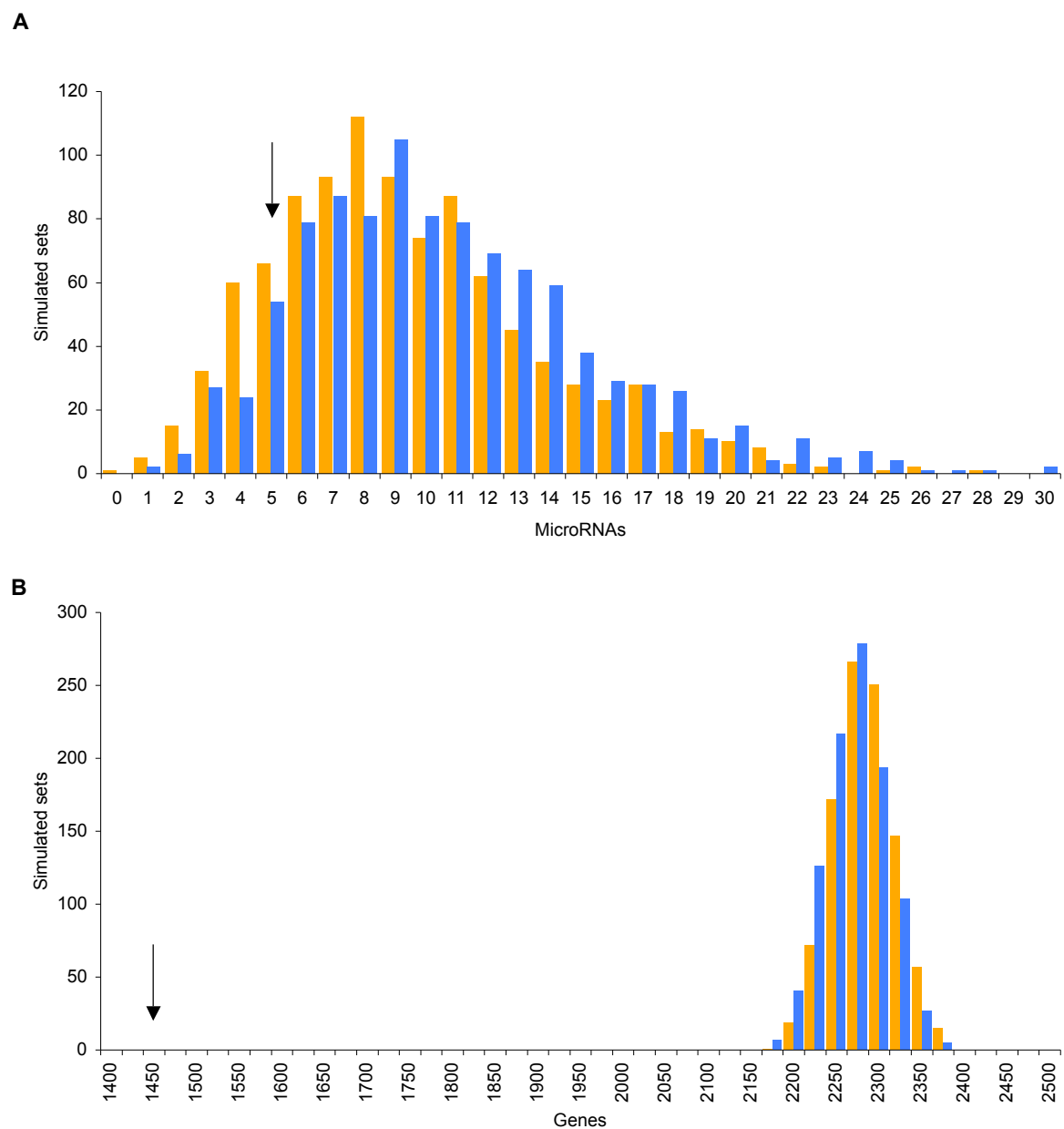


Figure S2. MicroRNAs and genes in repeat-poor SINE-free regions
Distributions of microRNA (A) and gene densities (B) in simulated sets of human SINE-free regions.
Only SINE-free regions with less than 50% masked sequence were considered, both in real and in simulated sets.
Arrows indicate the observed microRNA and gene numbers in the real data.